

123), lymphotoxin  $\alpha$  (LT $\alpha$ , SEQ ID NO 124), lymphotoxin  $\beta$  (LT $\beta$ , SEQ ID NO 125), FAS (SEQ ID NO 126), and CD40-L (ligand, SEQ ID NO 127), which are all members of the TNF family of proteins. One dot over a column indicates conservation across all sequences. Two dots over a column indicates the G(x)Y (glycine and tyrosine residues) identified in all the TNF related proteins.

Insert the following paragraphs on page 5, line 33:

SEQ ID NO 123 shows a portion of a human tumor necrosis factor protein.

SEQ ID NO 124 shows a portion of a human lymphotoxin  $\alpha$  protein.

SEQ ID NO 125 shows a portion of a human lymphotoxin  $\beta$  protein.

SEQ ID NO 126 shows a portion of a human FAS protein.

SEQ ID NO 127 shows a portion of a human CD40-L protein.

Replace the paragraph on page 20, lines 9-14, with the following:

The EDA1-II cDNA (SEQ ID NO 1) encodes a 391 residue protein (SEQ ID NO 2), 256 amino acids of which are encoded by new exons. EDA1-II is 94% identical to Tabby (SEQ ID NO 4), and includes a collagen-like domain with 19 repeats of a Gly-X-Y motif, interrupted by two amino acids between repeats 11 and 12 (FIG. 1). The full-length EDA1-II transcript can include a longer 3' UTR, because no polyadenylation signal sequence was identified, and Northern analysis indicates the transcript is 5-6 kb in length. The *Tabby* transcript is also 5-6 kb in length and has a 3' UTR of approximately 3.5 kb.

#### Remarks

The specification has been amended to include the sequence (SEQ ID NO) identifiers as requested by the Examiner. In addition, the sequence listing has been amended to include those sequences shown in figure 4 (SEQ ID NOS 123-127).

No new matter is added by this amendment.